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GenCore version 4.5
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SUMMARIES

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Signal; Calcium-binding; EGF-like domain; Coated pits;
Alternative splicing; POTENTIAL.
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01-CCT-1996 (Rel. 34, Lak: Esquence update)
01-CCT-1996 (Rel. 40, Last annotation update)
LOW-DENSITY LIFOPROTEIN RECEPTOR-RELATED PROFEIN 1 PRECURSOR (LRP)
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Seeppor; Transcription regulation; DNA-binding; Nuclear protein;
Zinc-finger.
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STRAIN-WHITE LEGHORN; TISSUE-Liver, and Ovary;
                                                                                                                                     30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
NUCLEAR HORMONE RECEPTOR FAMILY MEMBER NIR-25
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Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata;
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MAILTRE-5233191) Lubsed-7534747; Librar K., Ubbras L., Sermedis L.,
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Ferrorical Cit be seen (CHP) condition for the human popula.

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"Stratey to sequence the 80 score of the human Liel, gene coding for the Lipoprobean receptor related procedu. Receptor related procedu. Receptor related procedu. Receptor related procedu. Receptor stated procedu. Receptor States 44(1998).
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Sottrup-Jensen L.;
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Proteolytic processing of the 600 kd low density lipoprotein
receptor-related protein (LRP) occurs in a trans-Golgi
compartment.;
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Pfam; PF00057; 1dl_recept_a; 31.
Pfam; PF00058; 1dl_recept_b; 33.
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EMBL: AF058427; AAC64265.1; -.
PDB; 1CRB; 06-JAN-99.
EMBO J. 7:4119-4127(1988).
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		DBOXVI.: 3			1: 27.		Repeat; Endocytosis; Glycoprotein;	ike domain; Coated	u mondona	DECRETAIN 1	EXTRACELLIA (DOMENTAL)		POTENT	LDL-RECEPTOR CLASS A 1.	CLASS A 2		EGF-LIKE 2, CALCIUM-BINDING (POTENTIAL).	EGF-LIKE 3.		CLASS A 3	CLASS A 4	LDL-RECEPTOR CLASS A 5.	CLASS A B	× 0	CLASS A B	CLASS A S	THE CLASS A 1	EGF-LIKE 5.	DOS TAKE 3	DOD TING O	DOS-LINE O	EGF-17KE 10.	CLASS A 1	CLASS A 12	CLASS A 13	CLASS A 14	CLASS A 15	LDL-RECEPTOR CLASS A 16.	CLASS A 1/	RECEPTOR CLASS A 18	CLASS A 19	TKP 11	IX	JKE 13.	-RECEPTOR CLASS A 2	A 2	CLASS A 2	CLASS A 2	LDL-RECEPTOR CLASS A 29.	ECEPTOR CLASS A 3	CLASS A 3	SOUTHING 14.	2000-1100 16			EGF-LIKE 19.	EGP-LIKE 20.						
F_CA;	Ma; 31	ASX HV	BGF 1;	EGF_2; 8.	LDLRA	LDLRA	mbrane;	inding:	212	8 0 0	6177	4444	4544	99	110	149	189	250	843	768	933	5/5	1013	1000	1033	7707	787	7777	7071	1001	2105	2518	2563	2602	2641	2690	2732	2771	2814	0000	2000	2081	3000	3331	3371	3410	3450	3491	3533	3572	3611	3649	3695	3/33	3778	2061	7007	4232	4268	4304	4340	4375	
SM000179; EG SM00001: EG	100192; II	DS00010:	PS00022;	PS01186;	PS01209;	PS50068;	Transmen	aterum-Di	7 6	80			4445			111	150	474	803	708	668	200	100	1060	1100	77.07	1300	1183	1623	9701	2155	2478	2522	2564	2603	2642	2694	2732	27/72	2000	2000	2941	2002	3290	3332	3372	3411	3451	3492	3534	3573	3611	3652	3593	3/39	3034	4143	4196	4232	4268	4304	4340	
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WHITH   WEATH   12   12   CLUDERS R   R   R   R   R   R   R   R   R   R	Opery Match 10.0%; Score 26; DB 3; Length 58; Nest Local Similarity 23.5%; Pred. No. 33; Netches 4; Conservative 0; Mismatches 13; Indels 0; Gapes No. 2 CYNEWYNYNYNYN 18		55	TO 1.NEW-1.297 (New 1.3.7. Late Construence update) TO 1.SEV-1.297 (New 1.3.7. Late Annotation update) TO 1.SEV-1.298 (New 1.3.7. Late Annotation update) GO FORTINGWIN (NEW) GO FORTINGWIN (NEW 1.2.7. LATE ANNOTATION Update) GO RAMAGONE MEASORA (NEW 1.2. READING ANNOTATION Update) CO RAMAGONETRON, RECEIVED (NEW 1.2. DECOMPARIA READINGS GO REPRESENTED (NEW 1.2. DECOMPARIA READINGS				W. Metal-linding Petal-Lindings (Desiret; Chellation; Chellation
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Company of the compan	CLUSTER A. CLUSTER A. T. 176RBARFG0A32V9G CRCG4;	8; Score 26; DB 1; Length 58; 8; Pred. No. 33; 0; Mismatches 13; Indels 0; Gaps		per; 59 AA. sated) L sandoration update)	odnistrant brak : 80) Noverbrus skrouts). Demografia : Pleoryemata: Aktarides; takus:	9921011; "LL, Meejrup P., Depledge M.H., Knudsen J.; crospod crustacean mechallothionains with water and semi-terrestrial species."; 1995).	CYSTELIAS SAVEA A HIGH CONTENTO CYSTELIAS CHESTALES AND A HIGH CONTENTO CYSTELIAS CHESTALIAS CHESTALIAS INTERNATION CHESTALIAS CHESTALIAS INTERNATION CHESTALIAS CHES	Hilbhion.  Lithing Const.  Jack Shirth Condition.  Jack Shirth Condition.  CLISTER B.  CLISTER B.
Page	53 55 56 8; 5975 MW;	8; Score 26; DB 1; Length 58; 8; Pred. No. 33; 0; Mismatches 13; Indels 0; Gaps	2 CXXXXXXXXXXXXXXX 18 2 CXXXXXXXXXXXXXX 18 33 CTSGCXCPSKXXXXXX 49	h <sub>o</sub>	The Application of the Applicati	PARENISHES-970757; puembed-821011; Parenishes S.W.: Rederesm K.L.: Heolgraph C. Depledge M.H.: Krudsen J.; Perchalt General of F. Carley and C. Carley C. Ca	1 INVESTIGATION STRANGTONERS NAVE A HOST OWNERS OF CESSELER. RESIDENT STRAIN ST	SISTEP, POSSA 11MD.  Linearrow, TRADOLAGY, MARITHANIA, LANGERS, MARITHANIA, LANGERS, MARITHANIA, LANGERS, MARITHANIA, LANGERS, MARITHANIA, LANGERS, MARITHANIA, MA

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Tabuchi H., Lesslauer N.; Lahm H.-W., dentz R., det rumer ner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A., AND SEQUENCE OF 41-53; 110-124 AND 199-201
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EMBO J. 9:3269-3278(1990).
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TUMOR NECROSIS FACTOR BINDING PROTEIN 1.
                                                                                                                                                                                                                                                                                                        SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
TOMALIN: "HE DOMAIN THAT INDUCES A-SMASE IS PROGRELY IDENTICAL.
THE DEATH DOMAIN. THE N-SWASE ACTIVATION DOMAIN (NSD) IS BOTH
WEDESSARY AND SUPFICERN FOR ACTIVATION OF N-SWASE.
                                                                                                                                                                                                                                                                                                                                                                                                     --- SIMILARITY: CONTAINS A LA NERFAYRET-TYPE CYSTEINE-RICH REGION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00652; TWFR_NGFR_1; 3. PROSITE; PS50050; TWFR_NGFR_2; 3. PROSITE; PS50017; DEATH_DOMAIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL, M75866, AAA61201.1;
EMBL, M75864; AAA61201.1; JOINED.
EMBL, M75865, AAA61201.1; JOINED.
EMBL, M60275; AAA61201.1; JOINED.
EMBL, A21522; CAA01558.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00531; death; 1.
Pfam; PF00020; TNFR_C6; 4.
ProDom; PD000771; TNFR_C6; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; X55313; CAA39021.1; -. EMBL; M33394; AAA03210.1; -. EMBL; M5286; AAA36753.1; -. EMBL; M5866; AAA61201.1; -. EMBL; M75866; AAA61201.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SMART; SM00005; DEATH; J
SMART; SM00208; TNFR; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIR, A4899; GGHTI.
PIR, A35010, A35010.
PIR, S12057; S12057;
PIR, A8208; A38208.
PDB; ITNR; 31-JUL-94.
PDB; INCF; 07-DEC-95.
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Wandemburgh K., Goussin P., Nishharra D., Swidersful R., Silvestri G.,
Schordeste D. P., Silvestri G.S., Bird A.C., Sheffleld V.C., Silvestri G.,
Schordeste D. P., S.
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X. Toda T., Okti K., Makamara Y.;
Structure and chromosomal assignment of the human S1-5 gene (FBNL)
that is highly AmenOsques to fibrillin.*;
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Yn overspressed geder trascript in sensecut and quiescent I
fluodlasts exceding a novel protein in the epidermal growth
like repeat family stimates DNM synthesis.';
Mont. Cell. Biol., 15:120-128(1955).
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MEDLINE-95097983; PubMed-7799918;
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Score 26, DB 1, Length 493;
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TWRDIIRPS-97415782; pubbed-9268694;
Ozaki T, Kondo K., Nakamura Y., Ichiniya S., Nakagawara A.,
"Interaction of DAAI, a DAN-binding protein, with the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EGF-CONTAINING PIBULIN-LIKE EXTRACELLULAR
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20-ANG-2001 (Rel. 40, Last sequence update)
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21-ANG-2001 (Nel. 40, Last ammodation update)
21-ANG-2001 (Nel. 40, Taff Portson 2014)
21-ANG-2001 (Place 2014) (Taff Portson 2014)
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Bukaryota: Wetazoa: Chordata: Craniata: Vertebrata: Euteleostomi;
Mammalia: Euteria: Rodentia: Schuroquathi; Muridae; Murinaer Rattus.
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Bacchem, Biophys, Res. Commun. 237:245-250(1997).
-!- SUBCELLUIAR LOCATION: SECRETED.
-!- SIMILARITY: CONTAINS 6 EGF-LIKE DOMAINS.
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